

#5

SEQUENCE LISTING

<110> GREEN, Sol Alexander  
FRIEL, Ellen Nicola  
BEUNING, Lesley Leah  
MACRAE, Elspeth Ann

<120> Plant alpha farnesene synthase and polynucleotides encoding same

<130> 38-05

<140> 10/531,357

<141> 2005-04-14

<150> PCT/NZ2003/000229

<151> 2003-10-15

<150> NZ 521984

<151> 2002-10-15

<160> 14

<170> PatentIn version 3.1

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<212> DNA

<213> Malus domestica

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Arg Arg Ser Ala Asn Tyr Lys Pro Asn Ile Trp Lys Asn Asp Phe Leu  
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Asp Gln Ser Leu Ile Ser Lys Tyr Asp Gly Asp Glu Tyr Arg Lys Leu  
 50 55 60

Ser Glu Lys Leu Ile Glu Glu Val Lys Ile Tyr Ile Ser Ala Glu Thr  
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Met Asp Leu Val Ala Lys Leu Glu Leu Ile Asp Ser Val Arg Lys Leu  
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Gly Leu Ala Asn Leu Phe Glu Lys Glu Ile Lys Glu Ala Leu Asp Ser  
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Ile Ala Ala Ile Glu Ser Asp Asn Leu Gly Thr Arg Asp Asp Leu Tyr  
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Gly Thr Ala Leu His Phe Lys Ile Leu Arg Gln His Gly Tyr Lys Val  
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Asn His His Phe Ala His Leu Lys Gly Met Leu Glu Leu Phe Glu Ala  
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Ser Asn Leu Gly Phe Glu Gly Glu Asp Ile Leu Asp Glu Ala Lys Ala  
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Ser Leu Thr Leu Ala Leu Arg Asp Ser Gly His Ile Cys Tyr Pro Asp  
195 200 205

Ser Asn Leu Ser Arg Asp Val Val His Ser Leu Glu Leu Pro Ser His  
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Lys Asp Ile Cys Arg Val Asn Ala Thr Leu Leu Glu Leu Ala Lys Leu  
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Asn Phe Asn Val Val Gln Ala Gln Leu Gln Lys Asn Leu Arg Glu Ala  
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Ser Arg Trp Trp Ala Asn Leu Gly Phe Ala Asp Asn Leu Lys Phe Ala  
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Arg Asp Arg Leu Val Glu Cys Phe Ser Cys Ala Val Gly Val Ala Phe  
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Glu Pro Glu His Ser Ser Phe Arg Ile Cys Leu Thr Lys Val Ile Asn  
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Glu Leu Lys His Phe Thr Asn Ala Val Asp Arg Trp Asp Ser Arg Glu  
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Thr Glu Gln Leu Pro Glu Cys Met Lys Met Cys Phe Gln Val Leu Tyr  
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Met Ala Asp Phe Leu His Lys Asn Glu Asp Leu Leu Tyr Asn Ile Ser  
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Leu Ile Val Arg Leu Asn Asn Asp Leu Gly Thr Ser Ala Ala Glu Gln  
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Glu Arg Gly Asp Ser Pro Ser Ser Ile Val Cys Tyr Met Arg Glu Val  
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Asn Ala Ser Glu Glu Thr Ala Arg Lys Asn Ile Lys Gly Met Ile Asp  
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Asn Ala Trp Lys Lys Val Asn Gly Lys Cys Phe Thr Thr Asn Gln Val  
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Pro Phe Leu Ser Ser Phe Met Asn Asn Ala Thr Asn Met Ala Arg Val  
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acgatctcta tgctactgca ttacacttca agatcctcag gcagcatggc tataaagttt 180  
cacaagatat atttggtaga ttcattggatg aaaagggcac attagagaac caccatttcg 240  
cgcatttaaa aggaatgctg gaacttttcg aggcctcaaa cctgggtttc gaaggtgaag 300  
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Ser Asp Asn Leu Gly Thr Arg Asp Asp Leu Tyr Ala Thr Ala Leu His  
35 40 45

Phe Lys Ile Leu Arg Gln His Gly Tyr Lys Val Ser Gln Asp Ile Phe  
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Gly Arg Phe Met Asp Glu Lys Gly Thr Leu Glu Asn His His Phe Ala  
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His Leu Lys Gly Met Leu Glu Leu Phe Glu Ala Ser Asn Leu Gly Phe  
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Glu Gly Glu Asp Ile Leu Asp Glu Ala Lys Ala Ser Leu Thr Leu Ala  
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Leu Arg Asp Ser Gly His Ile Cys Tyr Pro Asp Ser Asn Leu Ser Arg  
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Asp Val Val His Ser Leu Glu Leu Pro Ser His Arg Arg Val Gln Trp  
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Phe Asp Val Lys Trp Gln Ile Asp Ala Tyr Glu Lys Asp Ile Cys Arg  
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Val Asn Ala Thr Leu Leu Glu Leu Ala Lys Leu Asn Phe Asn Val Val  
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Gln Ala Gln Leu Gln Lys Asn Leu Arg Glu Ala Ser Arg Trp Trp Ala  
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Asn Leu Gly Ile Ala Asp Asn Leu Lys Phe Ala Arg Asp Arg Leu Val  
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Glu Cys Phe Ala Cys Ala Val Gly Val Ala Phe Glu Pro Glu His Ser  
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